



Anforderung für molekularpathologische Diagnostik

Patient / Name	Patient / Geb.-Datum	Material / Materialnummer
anfordernder Arzt	Tel. (Rücksprache)	anfordernde Abteilung
Entität (Pflichtfeld)		Diagnose (Pflichtfeld)
FOKUS Targets / Gene / Fusionen / Komplettes Gen-Panel (bitte immer angeben)		
Anmerkungen		
Hinweise zur Abrechnung		Datum Probeneingang (Eintrag Institut für Pathologie)

Diagnostik nach Entitäten

Die Mutationsanalyse mit Next Generation Sequencing (NGS) erfolgt nach den Leitlinienempfehlungen zur molekularen Tumordiagnostik.

*NGS siehe Genliste **DNA/RNA-Panel** für detaillierte Informationen der untersuchten Genbereiche [siehe Rückseite](#)

Lungenkarzinom <input type="checkbox"/> Mutationsanalyse an FFPE (NGS DNA Panel2) <input type="checkbox"/> Mutationsanalyse an Liquid Biopsy (NGS DNA Panel1) <input type="checkbox"/> RNA-NGS (Genfusionen/Translokationen/METex 14-Skipping-Varianten)(NGS RNA Panel1) <input type="checkbox"/> FISH (ALK) <input type="checkbox"/> FISH (RET) <input type="checkbox"/> FISH (ROS1) <input type="checkbox"/> FISH (NTRK1-3) <input type="checkbox"/> FISH (MET)	Kolorektales Karzinom <input type="checkbox"/> Mutationsanalyse an FFPE (NGS DNA Panel3) <input type="checkbox"/> RNA-NGS (Genfusionen/Translokationen) (NGS RNA Panel1) HNPCC/LYNCH- Syndrom <input type="checkbox"/> MSI (Mikrosatelliteninstabilität mit Fragmentlängenanalyse) <input type="checkbox"/> MLH1, MSH2, MSH6, PMS2 MMR-Protein-Expression (IHC) <input type="checkbox"/> MLH1 Methylierung <input type="checkbox"/> BRAF (Mutationsanalyse an FFPE (NGS DNA Panel1)
Magenkarzinom <input type="checkbox"/> Mutationsanalyse an FFPE (NGS DNA Panel2) <input type="checkbox"/> FISH (HER2 Amplifikation)	Malignes Melanom/Aderhautmelanom <input type="checkbox"/> Mutationsanalyse an FFPE (NGS DNA Panel1) <input type="checkbox"/> Mutationsanalyse an Liquid Biopsy (NGS DNA Panel1) <input type="checkbox"/> FISH (HER2) <input type="checkbox"/> BRAF V600 Fast-Track
Neurologische Tumoren <input type="checkbox"/> Mutationsanalyse an FFPE (NGS DNA Panel5) <input type="checkbox"/> FISH (1p19q-Kodeletion) <input type="checkbox"/> FISH (MET) <input type="checkbox"/> FISH (CDKN2A) <input type="checkbox"/> MGMT Methylierung	Sarkom <input type="checkbox"/> Mutationsanalyse an FFPE (NGS DNA Panel1) <input type="checkbox"/> RNA - NGS (NGS RNA Panel2) <input type="checkbox"/> FISH (FUS/c16) Myxoid Sarkome <input type="checkbox"/> FISH (MDM2/c12) Liposarkom <input type="checkbox"/> FISH (SS18/c18) Synovialsarkom <input type="checkbox"/> FISH (EWSR1/c22) Ewing-Sarkom
Nierenzellkarzinom <input type="checkbox"/> FISH (TFE3)	Schilddrüsenkarzinom <input type="checkbox"/> Mutationsanalyse an FFPE (NGS DNA Panel1)
Urologische Tumore /Prostata-, Mamma-, Ovarial- und Peritonealkarzinom <input type="checkbox"/> Mutationsanalyse an FFPE (NGS DNA Panel1) <input type="checkbox"/> RNA-NGS (NGS RNA Panel1) <input type="checkbox"/> FISH (HER2)	Gallengangskarzinom <input type="checkbox"/> Mutationsanalyse an FFPE (NGS DNA Panel1) <input type="checkbox"/> RNA - NGS (NGS RNA Panel1)



Gastrointestinale Stromatumoren (GIST) <input type="checkbox"/> Mutationsanalyse an FFPE (NGS DNA Panel7)		Aneurysmatische Knochenzyste, Noduläre Fasciitis <input type="checkbox"/> FISH (USP6/c17)	
Pankreaskarzinom <input type="checkbox"/> Mutationsanalyse an FFPE (NGS DNA Panel1 und/oder 6) <input type="checkbox"/> MSI (Mikrosatelliteninstabilität)		Myeloische Leukämie (AML), MDS, Myelofibrose, Myeloproliferative Neoplasie <input type="checkbox"/> Mutationsanalyse (NGS DNA Panel4)	
Erregerdiagnostik <input type="checkbox"/> B. burgdorferi <input type="checkbox"/> Humanes Papilloma Virus (HPV) <input type="checkbox"/> M. Whipple <input type="checkbox"/> Fungi <input type="checkbox"/> TB (Mycobakterium tuberculosis) <input type="checkbox"/> Herpes Viren (CMV, HSV, HHV6, EBV, VZV) <input type="checkbox"/> MOTT (Mycobacteria other than TB)		Lymphomdiagnostik <input type="checkbox"/> B-Zell-Klonalitätsanalyse (IgH, IgK, IgL) <input type="checkbox"/> T-Zell-Klonalitätsanalyse (TCR β , TCR γ) <input type="checkbox"/> FISH BCL2::IgH / t(14;18) PCR	
FISH-Analysen (Amplifikationen/Fusionen/Translokationen) <input type="checkbox"/> 1p19q <input type="checkbox"/> BCL2 (BreakApart) <input type="checkbox"/> EWSR1 (BreakApart) <input type="checkbox"/> HER2 Mamma/Magen (Amplifikation) <input type="checkbox"/> MYC (BreakApart) <input type="checkbox"/> HER2 sonstiges (Amplifikation) <input type="checkbox"/> MET (Amplifikation)		<input type="checkbox"/> NTRK1 (BreakApart) <input type="checkbox"/> NTRK2 (BreakApart) <input type="checkbox"/> NTRK3 (BreakApart) <input type="checkbox"/> RET (BreakApart) <input type="checkbox"/> TFE3 (BreakApart) <input type="checkbox"/> BCL::IgH t (14;18) (Fusion) <input type="checkbox"/> CDKN2A (homozygote Deletion)	
Sonstiges <input type="checkbox"/> DNA-Extraktion <input type="checkbox"/> RNA-Extraktion		<input type="checkbox"/> MGMT-Methylierung <input type="checkbox"/> MLH1-Methylierung <input type="checkbox"/> MSI	
Weitere DNA-NGS (Mutationen, InDels, CNVs), RNA-NGS (Fusionen/Rearrangements/Translokationen/METex14Skipping) (Targets bitte angeben) <input type="checkbox"/>			



*NGS Genliste DNA/RNA-Panel

* NGS Panel Genlisten:

DNA Panel 1: "mini-nNGM" (DNA-NGS): ALK, BRAF, EGFR, KRAS, MET, NRAS, PIK3CA

DNA Panel 2: "nNGM Panel" (DNA-NGS): ALK, BRAF, CTNNB1, EGFR, ERBB2, FGFR1-4, HRAS, IDH1, IDH2, KEAP1, KRAS, MAP2K1, MET, NRAS, NTRK1-3, PIK3CA, PTEN, RET, ROS1, STK11, TP53, TRDMT1

DNA Panel 3: "Kolon-Panel" (DNA-NGS): APC, BRAF, CTNNB1, FBXW7, KRAS, NRAS, PIK3CA, SMAD4, TP53

DNA Panel 4: "Myeloid-Onko-Panel" (DNA-NGS): ANKRD26, ASXL1, BCOR, BCORL1, BRAF, CALR, CBL, CEBPA, CSF3R, CXCR4, DDX41, DNMT3A, ETKN1, ETV6, EZH2, FLT3, GATA2, GNB1, IDH1, IDH2, JAK2, KIT, KRAS, MPL, MYD88, NF1, NOTCH1, NPM1, NRAS, PHF6, PIGA, PPM1D, PRPF8, PTPN11, RAD21, RUNX1, SF3B1, STAG2, SRSF2, TERC, TERT, TET2, TP53, U2AF1, WT1, ZRSR1

DNA Panel 5: "Neuro-Onko-Panel" (DNA-NGS): AKT1, ATRX, BRAF, CDKN2A, CIC, DAXX, EGFR, GNA11, GNAQ, H3F3A, H3F3B, IDH1, IDH2, KDM6A, KLF4, KLLN, NF1, NF2, PIK3CA, PIK3R1, POLR2A, PTEN, SMARCB1, SMO, STAG2, SUFU, TERT Promotor, TP53, TRAF7

DNA Panel 6: "BRCA" (DNA-NGS): BRCA1 und BRCA2

DNA Panel 7: "AIT" (Actionable Insights Tumor) (DNA-NGS): AKT1, ALK, BRAF, CTNNB1, EGFR, ERBB2, ERBB3, ESR1, FOXL2, GNA11, GNAQ, IDH1, IDH2, KIT, KRAS, MET, NRAS, PDGFRA, PIK3CA, RAF1, RET, TP53

RNA Panel 1: "Fusionen nach nNGM" (RNA-NGS): ALK, BAG4, BAIAP2L1, BRAF, CCDC6, CD74, COPA, CIAO1, CUX1, EGFR, EML4, ERC1, ETV6, EZR, FGFR1, FGFR2, FGFR3, FGFR4, GOPC, HIP1, KIF5B, KLC1, LRIG3, MET, MET, (Exon-14-Skipping) MPRIP, MRPS14, NRG1, NTRK1, NTRK2, NTRK3, NUTM1, RAD51, RET, ROS1, SDC4, SLC34A2, STRN, TACC3, TFG, TPR, TRIM33, UBE3C

RNA Panel 2: "Sarkom NGS Panel" (RNA-NGS): ALK, BCOR, BRAF, CAMTA1, CCNB3, CIC, EPC1, EWSR1, FOSB, FOXO1, FUS, GLI1, HMGA2, JAZF1, MEAF6, MKL2, NTRK1, NTRK2, NTRK3, PDGFB, PLAG1, ROS1, SS18, STAT6, TAF15, TCF12, TFE3, TFG, USP6, YWHAE

nNGM: Nationales Netzwerk Genomische Medizin Lungenkrebs; www.nngm.de